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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/902,188A

DATE: 12/06/2001 TIME: 11:36:38

Input Set : N:\Crf3\RULE60\09902188A.txt
Output Set: N:\CRF3\12062001\I902188A.raw

SEQUENCE LISTING

```
3 (1) GENERAL INFORMATION:
      5
             (i) APPLICANT: Bisgard-Frantzen, Henrik
                            Svendsen, Allan
      6
                                                                     ENTERED
      7
                            Borchert, Torben Vedel
            (ii) TITLE OF INVENTION: AMYLASE VARIANTS
      9
     11
           (iii) NUMBER OF SEQUENCES: 32
            (iv) CORRESPONDENCE ADDRESS:
     13
                  (A) ADDRESSEE: Novo Nordisk of North America, Inc.
     14
     15
                  (B) STREET: 405 Lexington Avenue, Suite 6400
     16
                  (C) CITY: New York
     17
                  (D) STATE: New York
     18
                  (E) COUNTRY: U.S.A.
     19
                  (F) ZIP: 10174-6401
     21
             (v) COMPUTER READABLE FORM:
     22
                  (A) MEDIUM TYPE: Floppy disk
     23
                  (B) COMPUTER: IBM PC compatible
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     24
     25
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
     27
            (vi) CURRENT APPLICATION DATA:
C--> 28
                  (A) APPLICATION NUMBER: US/09/902,188A
C--> 29
                  (B) FILING DATE: 10-Jul-2001
     30
                  (C) CLASSIFICATION:
     32
           (vii) PRIOR APPLICATION DATA:
     33
                  (A) APPLICATION NUMBER: 09/354,191
     34
                  (B) FILING DATE:
     36
          (viii) ATTORNEY/AGENT INFORMATION:
     37
                  (A) NAME: Lambiris, Elias J.
     38
                  (B) REGISTRATION NUMBER: 33,728
     39
                  (C) REFERENCE/DOCKET NUMBER: 4318.204-US
     41
            (ix) TELECOMMUNICATION INFORMATION:
     42
                  (A) TELEPHONE: 212 867 0123
     43
                  (B) TELEFAX: 212 867 0298
        (2) INFORMATION FOR SEQ ID NO: 1:
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     48
             (i) SEQUENCE CHARACTERISTICS:
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     49
     50
                  (B) TYPE: amino acid
     51
                  (C) STRANDEDNESS: single
     52
                  (D) TOPOLOGY: linear
     54
            (ii) MOLECULE TYPE: peptide
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             His His Asn Gly Thr Asn Gly Thr Met Met Gln Tyr Phe Glu Trp Tyr
     59
                                                  10
     61
             Leu Pro Asn Asp Gly Asn His Trp Asn Arg Leu Arg Asp Asp Ala Ala
     62
                         20
                                              25
     64
             Asn Leu Lys Ser Lys Gly Ile Thr Ala Val Trp Ile Pro Pro Ala Trp
     65
                     35
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Input Set : N:\Crf3\RULE60\09902188A.txt
Output Set: N:\CRF3\12062001\I902188A.raw

67 68	Lys	Gly 50	Thr	Ser	Gln	Asn	Asp 55	Val	Gly	Tyr	Gly	Ala 60	Tyr	Asp	Leu	Tyr
70 71	Asp 65	Leu	Gly	Glu	Phe	Asn 70	Gln	Lys	Gly	Thr	Val 75	Arg	Thr	Lys	Tyr	Gly 80
73 74	Thr	Arg	Asn	Gln	Leu 85	Gln	Ala	Ala	Val	Thr 90	Ser	Leu	Lys	Asn	Asn 95	Gly
76 77	Ile	Gln	Val	Tyr 100	Gly	Asp	Val	Val	Met 105	Asn	His	Lys	Gly	Gly 110	Ala	Asp
79 80	Gly	Thr	Glu 115	Ile	Val	Asn	Ala	Val 120	Glu	Val	Asn	Arg	Ser 125	Asn	Arg	Asn
82 83	Gln	Glu 130	Thr	Ser	Gly	Glu	Tyr 135	Ala	Ile	Glu	Ala	Trp 140	Thr	Lys	Phe	Asp
85 86	Phe 145	Pro	Gly	Arg	Gly	Asn 150	Asn	His	Ser	Ser	Phe 155	Lys	Trp	Arg	Trp	Tyr 160
88 89	His	Phe	Asp	Gly	Thr 165	Asp	Trp	Asp	Gln	Ser 170	Arg	Gln	Leu	Gln	Asn 175	Lys
91 92	Ile	Tyr	Lys	Phe 180	Arg	Gly	Thr	Gly	Lys 185	Ala	Trp	Asp	Trp	Glu 190	Val	Asp
94 95	Thr	Glu	Asn 195	Gly	Asn	Tyr	Asp	Tyr 200	Leu	Met	Tyr	Ala	Asp 205	Val	Asp	Met
97 98	Asp	His 210	Pro	Glu	Val	Ile	His 215	Glu	Leu	Arg	Asn	Trp 220	Gly	Val	Trp	Tyr
100 101	Thr 225		Thr	Leu	ı Asn	Leu 230	_	Gly	Phe	e Arg	11e	_	Ala	. Va]	L Lys	s His 240
103 104			Tyr	Ser	Phe 245	Thr		J Asp	Trp	Leu 250	Thr		Val	Arg	J Asi 25!	n Thr
106 107	Thr	Gly	Lys	Pro 260	Met		Ala	a Val	. Ala	Glu		Trp	Lys	Asr 270	ı Ası	p Leu
109 110	Gly	/ Ala	11e 275	Glu		туг	Let	1 Asn 280	Lys		Ser	Trp	Asn 285	His		r Val
112 113	Phe	Asp 290	Val		Leu	ı His	Ty:	Asn		ı Tyr	Asn	Ala	Ser		n Sei	r Gly
115 116	Gl ₃ 305	y Tyr		Asp	Met	Arg 310	Asr		. Leu	ı Asn	Gly 315	ser Ser		Va]	l Glı	Lys 320
118 119			Thr	His	325	val		Phe	val	Asp 330	Asn		Asp	Sei	Glr 33!	n Pro
121 122	Gly	g Glu	ı Ala	Leu 340	ı Glu		Phe	val val	. Glr 345	ı Gln		Phe	Lys	Pro 350) Le	ı Ala
124 125	Туг	Ala	Leu 355	Val		ı Thr	Arg	g Glu 360	Glr		Туг	Pro	Ser 365	Va]		e Tyr
127 128	Gly	7 Asp 370	туг		Gly	, Ile	Pro	Thr		s Gly	Val	. Prc 380	Ala		Lys	s Ser
130 131	Lys 385	: Ile		Pro	Leu	1 Leu 390	Glr		Arg	g Gln	Thr 395	Phe		Туз	Gly	y Thr
133			asp	туг		a Asp		s His	Asp		Ile		Trp	Thi		400 g Ģlu
134 136	Gly	y Asn	ser				Asr	n Ser				Thr	· Ile			r Asp
137 139	Gly	y Pro	Gly	420 Gly		Lys	Tr	Met	425 Tyr		Gly	Lys	Asn	430 Lys		a Gly

Input Set : N:\Crf3\RULE60\09902188A.txt
Output Set: N:\CRF3\12062001\1902188A.raw

140				435					440					445			
142		Gln	Val	Trp	Arg	Asp	Ile	Thr	Gly	Asn	Arg	Thr	Gly	Thr	Val	Thr	Ile
143			450	_	_	_		455					460				
145		Asn	Ala	Asp	Gly	Trp	Gly	Asn	Phe	Ser	Val	Asn	Gly	Gly	Ser	Val	Ser
146		465					470					475					480
148		Val	${\tt Trp}$	Val	Lys	Gln											
149						485						,					
152	(2)	INFO	RMAT	ION I	OR S	SEQ :	D NO): 2	:								
154		(i)					CERIS										
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156			(B) TYPE: amino acid														
157			(C) STRANDEDNESS: single														
158			(D) TOPOLOGY: linear														
160			MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 2:														
162												~ 1	_	m 1	a 1	m	
164			His	Asn	GLY	_	Asn	GLY	Thr	Met		GIn	Tyr	Phe	GIU		HIS
165		1	_	_	_	5	_		_	•	10	.	3	•	•	15	a
167		Leu	Pro	Asn	_	GIĀ	Asn	HIS	Trp		Arg	ьeu	Arg	Asp		Ата	Ser
168		3	T	3	20	3	61	T1_	m1	25	T1.	m	T1.	Dmo	30	71	m~~
170		ASN	ьeu	Arg	ASI	Arg	GIY	ше	40	Ald	ire	тгр	TTE	Pro 45	PIO	Ala	пр
171 173		T ***	C1		Cor	Cln	7 an	N a ra		Cly	Птт	C117	λla	Tyr	λαn	Lau	Пагт
174		гуу	50	TIII	ser	GIII	ASII	55	val	GLY	TYT	GIY	60	TYT	АБР	пеп	- Y -
176) cn		Clv	Glu	Dho	Aen		Luc	Glw	Thr	Va 1		Thr	T.v.c	ጥህጉ	Glv
177		65	пеп	GLY	GIU	rne	70	GIII	цуз	Gry	1111	75	пту	1111	ט עם	- 7 -	80
179			Arσ	Ser	Gln	Len		Ser	λla	Tle	His		Leu	Lys	Asn	Asn	
180			**** 9	001	01	85	OLU	001			90			-1-		95	0-1
182		Val	Gln	Val	Tvr		Asp	Val	Val	Met		His	Lvs	Gly	Gly	Ala	Asp
183					100	4				105			-	_	110		•
185		Ala	Thr	Glu	Asn	Val	Leu	Ala	Val	Glu	Val	Asn	Pro	Asn	Asn	Arg	Asn
186				115					120					125			
188		Ğln	Glu	Ile	Ser	Gly	Asp	Tyr	Thr	Ile	Glu	Ala	Trp	Thr	Lys	Phe	Asp
189			130					135					140				
191		Phe	Pro	Gly	Arg	Gly	Asn	Thr	Tyr	Ser	Asp	Phe	Lys	\mathtt{Trp}	Arg	\mathtt{Trp}	Tyr
192		145					150					155					160
194		His	Phe	Asp	Gly		Asp	Trp	Asp	Gln		Arg	Gln	Phe	Gln		Arg
195						165	_		_		170					175	
197		Ile	Tyr	Lys		Arg	Gly	Asp	Gly		Ala	Trp	Asp	Trp		Val	Asp
198				_	180	_	_	_		185		_	_ •	_	190	_	
200		Ser	Glu		GIĀ	Asn	Tyr	Asp		Leu	Met	Tyr	Ala	Asp	Val	Asp	Met
201		_		195	01	**- 1	**- 7		200	+	•	3	m	205	a 1	m	m
203		Asp		Pro	GIU	vaı	vaı		GIU	Leu	Arg	Arg		Gly	GIU	тгр	Tyr
204		mb se	210	mh m	т о	N a n	T 011	215	C1	Dho	λ ~~	тіс	220	71 -	Wa I	T 170	цiо
206			ASII	Tnr	ьeu	ASII		Asp	GTÅ	rne	HI.G	235	нар	Ala	val	пÃR	240
207 209		225	Two	Птт∽	S0~	Dhe	230	7~~	y c.r.	Пrr	Ten		Wic.	Val	λκα	λer	
210		тте	пур	тйг	3eT	245	THE	AI 9	vab	ттЪ	250	TIII	HIS	vaı	тту	255	пта
210		ጥኮኮ	Glv	Lare	Glu		Phe	בו∆	Val	Δla		Phe	Trn	Lys	Asn		Len
213		T11T	GIY	пур	260	rie t	FIIC	лта	Val	265	JIU	1110	112	כעם	270	ASP	cu
ويدم					200					200					2,0		

Input Set : N:\Crf3\RULE60\09902188A.txt
Output Set: N:\CRF3\12062001\I902188A.raw

215		Gly	Ala	Leu	Glu	Asn	Tyr	Leu		Lys	Thr	Asn	Trp		His	Ser	Val
216		Dha	3 am	275	Dwo	T 0.11	774 ~	M	280	T 011	m	7 ~ ~	71.	285	7	Com	C1
218		Phe	290	Val	PIO	ьeu	пıs	295	ASII	ьeu	TAL	ASII	300	ser	ASII	ser	СТА
219		C1		Птт	7 an	Wat	7.7.		T 011	T 0.11	7 a n	C1		17-1	17-1	C1 n	T ***
221 222		305	ASII	Tyr	ASP	Met	310	гуу	ьeu	ьeu	ASII	315	TIIT	val	vaı	GIII	320
224			Dro	Mot	uia	ת 1 ת		mh~	Dho	17-1	λου		цiс	7.00	cor	Cln	
224		urs	PIO	Met	птъ	325	Val	THI	Pile	val	330	ASII	птѕ	Asp	ser	335	PIO
		C1**	C1.,	Com	T 011		C07	Dho	Wa I	C1 n		Пхх	Dho	T	Dro		7 J ¬
227 228		GIY	GIU	Ser	340	GIU	261	Pne	val	345	GIU	тър	Pne	пур	350	ьeu	AId
230		Пага	אן א	Leu		Tau	Thr	λνα	Cl u		Cl v	Птт	Dro	Sar		Dho	Пагъ
231		TYT	АТа	355	110	пец	1111	ni 9	360	GIII	GLY	TYT	110	365	Val	riic	TYT
233		Cly	λan	Tyr	Пттъ	Cly	Tla	Dro		Wic	Sar	Wa l	Dro		Ma+	Lazo	λla
234		GIY	370	тут	TÄT	GLY	116	375	T111	nrs	261	Val	380	Ата	Mec	шу.ъ	Ата
234		Lvc		Asp	Dro	Tla	Lau		λla	λνα	Gl n	λcn		λla	Пагт	C1 17	Thr
237		385	116	нар	FIO	116	390	GIU	AIG	AIG	GIII	395	FILE	Ala	TYT	GIY	400
239			Hic	Asp	Фулт	Dhe		Hie	Hic	Δen	Tle		G) v	Фтъ	Πhr	Δτα	
240		GIII	1113	нар	- Y -	405	изр	1113	1113	non	410	110	GLY	111	1111	415	GIU
242		Glw	λen	Thr	Пhr		Dro	λen	Ser	Clv		λ 1 =	Фhr	Tle	Mo+		Δen
243		GLY	H211	1111	420	1113	110	ASII	SCI	425	пец	лта	1111	110	430	JCI	пор
245		Glv	Pro	Gly		Glu	T.v.c	Ψrn	Met		Val	Glv	Gln	Δen		Δla	Glv
246		Cly	110	435	OLY	OLU	Lys		440	+1-	V 41	011	0111	445	1170	1114	011
248		Gln	Val	Trp	Hic	Δsn	Tle	Thr		Δsn	T.vc	Pro	Glv		Va 1	Thr	Tle
249		GIII	450	115	1110	пор	110	455	O _T	11511	17.5	110	460	1111	•41		110
251		Δsn		Asp	Glv	Trn	Δla		Phe	Ser	Va l	Asn		Glv	Ser	Val	Ser
252		465			011		470		1	001	•	475	011		001		480
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255					-1-	485											
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260	\ - <i>/</i>			JENCE													
261		\ - <i>\</i>) LEI						3							
262				TYI													
263			-) STI					le								
264			•	, TOI				-									
266		(ii)	•	2													
268		(xi)				_	-		EQ II	ON C	: 3:						
270				Pro								Tyr	Phe	Glu	Trp	Tyr	Leu
271		1				5	_				10	_			_	15	
273		Pro	Asp	Asp	Gly	Thr	Leu	Trp	Thr	Lys	Val	Ala	Asn	Glu	Ala	Asn	Asn
274					20					25					30		
276		Leu	Ser	Ser	Leu	Gly	Ile	Thr	Ala	Leu	Trp	Leu	Pro	Pro	Ala	Tyr	Lys
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Input Set : N:\Crf3\RULE60\09902188A.txt Output Set: N:\CRF3\12062001\1902188A.raw

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115		Thr	Glu	Trp		Asp	Ala	Val	Glu		Asn	Pro	Ser	Asp	Arq	Asn	Gln
294 . Glu				_		•									_		
130		Glu	Ile	Ser	Glv	Thr	Tyr	Gln		Gln	Ala	Trp	Thr	Lys	Phe	Asp	Phe
Pro					-		•					-		-		-	
145		Pro	Gly	Arq	Gly	Asn	Thr	Tyr	Ser	Ser	Phe	Lys	Trp	Arg	Trp	Tyr	His
Solid			-	,	_			•					-	_	-	•	
301			Asp	Glv	۷al	Asp	Trp	Asp	Glu	Ser	Arq	Lvs	Leu	Ser	Arq	Ile	Tvr
1				1		_						_1 -					-
304 Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Asp Leu Hyr Asp Leu Hyr Asp Leu Hyr Leu Lyr Bull Asp Leu Lyr Bull Lyr Bull Hyr Lyr Bull Lyr Bull Lyr Lyr Tyr Tyr Tyr Tyr Lyr Lyr Tyr Tyr Tyr Tyr Lyr Ly		Lvs	Phe	Arg	Glv	Ile	Glv	Lvs	Ala	Trp	Asp	Trp	Glu	Val	Asp	Thr	Glu
Ash Sin Ash Sin Tyr Ash Tyr Leu Met Tyr Ala Ash Leu Ash Met Ash His 200		-1-		5	_			_1 -		_							
307 Pro GL 141 Val Val Val Val Thr Glu Leu Lys Ser Trp Gly Lys Trp Val Val Val San Asn 215 220 220 325 320 32		Asn	Glv	Asn		Asp	Tvr	Leu	Met	Tyr	Ala	Asp	Leu	Asp	Met	Asp	His
Pro Glu Val Val Thr Glu Leu Lys Ser Trp Gly Lys Trp Tyr Val Asn Silo 210 215 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 225 220 227 220 220 227 220 220 227 220			4		- 4	•	•			-		•				•	
310 210 Thr Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Phe Phe Phe Asp Asp Trp Leu Ser Asp Asp Gly 240 Asp Trp Gly 250 Trp 240 240 240 240 240 240 240 240 240 240 240 260 250 Trp 250 Trp 255 250 Trp 255 250 Trp 255 270 255 270 275 270 275 270 275 270 275 270 275 270 275 270 275 270 275 270 275 275 270 275 270 275 270 275 270 275 270 <td></td> <td>Pro</td> <td>Glu</td> <td></td> <td>Val</td> <td>Thr</td> <td>Glu</td> <td>Leu</td> <td>Lvs</td> <td>Ser</td> <td>Trp</td> <td>Glv</td> <td>Lvs</td> <td>Trp</td> <td>Tvr</td> <td>Val</td> <td>Asn</td>		Pro	Glu		Val	Thr	Glu	Leu	Lvs	Ser	Trp	Glv	Lvs	Trp	Tvr	Val	Asn
Thr Thr Asn Ile Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys 240	,								_1 -			2		•	-		
313		Thr		Asn	Ile	Asp	Glv		Ara	Leu	Asp	Ala	Val	Lvs	His	Ile	Lvs
Simple Ser Phe Phe Phe Phe Phe Asp Trp Leu Ser Asp Val Arg Ser Gln Thr Gly 245 255 318 Lys Pro Leu Phe Thr Val Gly Glu Tyr Trp Ser Tyr Asp Ile Asn Lys Lys 126 265 265 265 265 270 2									5					-1 -			
316			Ser	Phe	Phe	Pro		Trp	Leu	Ser	Asp		Ara	Ser	Gln	Thr	
318 Lys Pro Leu Phe Thr Val Gly Gly Tyr Tyr Asp Ile Asn Lys 260 265 327 327 327 327 321 Leu His Asn Tyr Ile Met Lys Thr Asn Gly Thr Asn Lys Ser Gly Gly Thr Asn Asn Lys Pro Thr Asn Thr Asn Asn Thr Pro Thr Asn Thr Asn Asn Thr Asn Asn Thr Asn Asn Asn Thr Asn Asn Asn Thr Asn Asn Asn <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td> F</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>1</td>							F										1
319 Leu His Asn Tyr 11e Met Lys Tile Met Lys Thr Asn Gly Thr Met Ser Leu Phe Asp 275 270 280 11e Met Ser Lys Ser Gly Gly Thr Met Ser Leu Phe Asp 285 11e Phe Asp 31e Phe Asp 31e 11e Phe Asp 31e <		Lvs	Pro	Leu	Phe		Val	Glv	Glu	Tvr		Ser	Tvr	Asp	Ile		Lvs
321 Leu His Asn Tyr 11e Met Lys Thr Asn Gly Thr Met Ser Leu Phe Asp 280 Asn 285 Leu Phe Asp 280 Asn 285 Leu Phe Asp 285 Asn 290 Asn 290 Leu Phe Asp 295 Thr Ala Ser Lys Ser Gly Gly Thr 300 Ser Gly Gly Thr 300 Thr 325 Asp 290 Asn 295 Thr Leu Asp 295 Thr Asn Thr Leu Asp 300 Met Lys Asp Gly Thr 300 Asp 300								1		_			-1-				-1-
322 Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser Lys Ser Gly Gly Thr 229 Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser Lys Ser Gly Gly Thr 325 Ser Gly Gly Thr 300 327 Phe Asp Met Asp Met Arg Thr Leu Met Thr Asn Thr Leu Met Thr Asn 315 Thr Leu Met Lys Asp Gln Pro 320 330 Thr Leu Ala Ser Thr Leu Met Thr Asn Thr Leu Met Thr Asn Thr Leu Met Thr Asn 315 Thr Glu Pro Gly Gln 320 330 Thr Leu Ala Ser Thr Phe Val Asp Ser Thr Phe Val Asp Ser Thr Ser Lys Thr Ser Lys Thr Asn 330 Thr Glu Pro Gly Gln 335 333 Ala Leu Gln Ser Thr Arg Gln Glu Gly Tyr Phe Lys Pro Cys Val Pro Ser Leu Ser Tyr Ala 340 340 336 Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly Asp 360 339 Tyr Tyr Gly Ile Pro Gln Tyr Asn Ile Pro Ser Leu Lys Ser Lys Ile 340 340 370 341 Tyr Tyr Gly Leu Ser Lys Tyr Asn Ile Pro Ser Leu Lys Ser Lys Ile 360 342 Asp Pro Leu Leu Ile Ile Gly Tyr Ala Tyr Ala Tyr Ala Tyr Ala Ser Lys Thr Gly Thr Gln His Asp 360 343 Asp Tyr Leu Asp His Ser Asp Ile Ile Gly Trp Thr Arg Glu Gly Val 400 345 Asp Tyr Leu Asp His Ser Lys Tyr Asp Ile Ile Gly Tyr Thr Arg Gly Thr Gly Val 410 346 Thr Glu Lys Fro Gly Ser Lys Tyr Asp Ile Ile Gly Tyr Thr Arg Gly Thr Asp Gly Val 410 348 Thr Glu Ser Lys Tyr May Gly Ser Lys Tyr May Gly Lys Gln His Asp Gly Val 445 351		Leu	His	Asn		Tle	Met	Lvs	Thr		Glv	Thr	Met	Ser		Phe	Asp
324 Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser Lys Ser Gly Gly Thr 290 295 Thr Ala Ser Lys Ser Gly Gly Thr 300 327 Phe Asp Met Arg Thr Leu Met Thr Asn Thr Leu Met Sun January Met Lys Asp Gln Pro 320 328 305 Thr Leu Ala Val Thr Phe Val Asp Asn His Asp Thr Glu Pro Gly Gln 331 310 Thr Leu Ala Val Thr Phe Val Asp Asn His Asp Thr Glu Pro Gly Gln 333 325 330 Thr Leu Gln Ser Trp Val Asp Pro Trp Phe Val Asp 345 330 Thr Glu Pro Gly Gln Asp 355 330 330 335 336 337 336 337 337 337 337		Lou			-1-			-1-			0-1						
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357 Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly Ser Val Ser Val Trp		Asp	Gly	Trp	Gly	Glu	Phe	Lys	Val	Asn	Gly	Gly	Ser	Val	Ser	Val	Trp
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360 Val Pro Arg Lys Thr Thr Val Ser Thr Ile Ala Trp Ser Ile Thr Thr			Pro	Arg	Lys	Thr	Thr	Val	Ser	Thr	Ile	Ala	Trp	Ser	Ile	Thr	Thr
361 485 490 495	361			_	-								_				

VERIFICATION SUMMARY

DATE: 12/06/2001

PATENT APPLICATION: US/09/902,188A

TIME: 11:36:40

Input Set : N:\Crf3\RULE60\09902188A.txt
Output Set: N:\CRF3\12062001\I902188A.raw

L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]